

Fig. 1.

α411-wr MYSLVFLVILMCIPIF-----SFOITVYDDKSVCI-DSDNKEYMGIEVYVEA 42
 ctkbp-11ster --MKQYIYLAACMLAAAMPASLQSSSSSCTEEENKHHMGIDYIILKV 48

α411-wr TLDE--PLRGTTCESKIHKYGA-----ESEDSTSVEDVDPTTYS 63
 ctkbp-11ster TTKQDQTPTNDKICQSVTEITETESDPEVESEDDSTSVEDVDPTTYS 98

α411-wr VSNGGLNISVDLNLNCFLNFTVGVYTNRDIVYAKFAISLDPWITTEPINSM 113
 ctkbp-11ster IIGGGLRMNFGEFTKC--POIKSIISESAAGNTVNARLSSVSPGGKOSPAT 147

α411-wr HDDLVKLTTEECIVDIYLYLKCCEVDKTKDFMKTNGNRLKPRDFKITVPPSNVGS 163
 ctkbp-11ster REEALAMIKDCQEVSIIRCSSEEKQSDIKIHPVLSNISHKKVSYEDIIG 197

α411-wr MIELOSQYCVNDVYTYVKIYDECGNICKQHSIPIILRQYFTTKNG-- 206
 ctkbp-11ster STIVDTK-CVKNLEFSVRIGDMC---KESSELEVKGDFKYVDGSA SEGAT 243

α411-wr -OPRKLK KFDNCI-
 ctkbp-11ster DOTSLIDSTKLKACV

Fig.2.

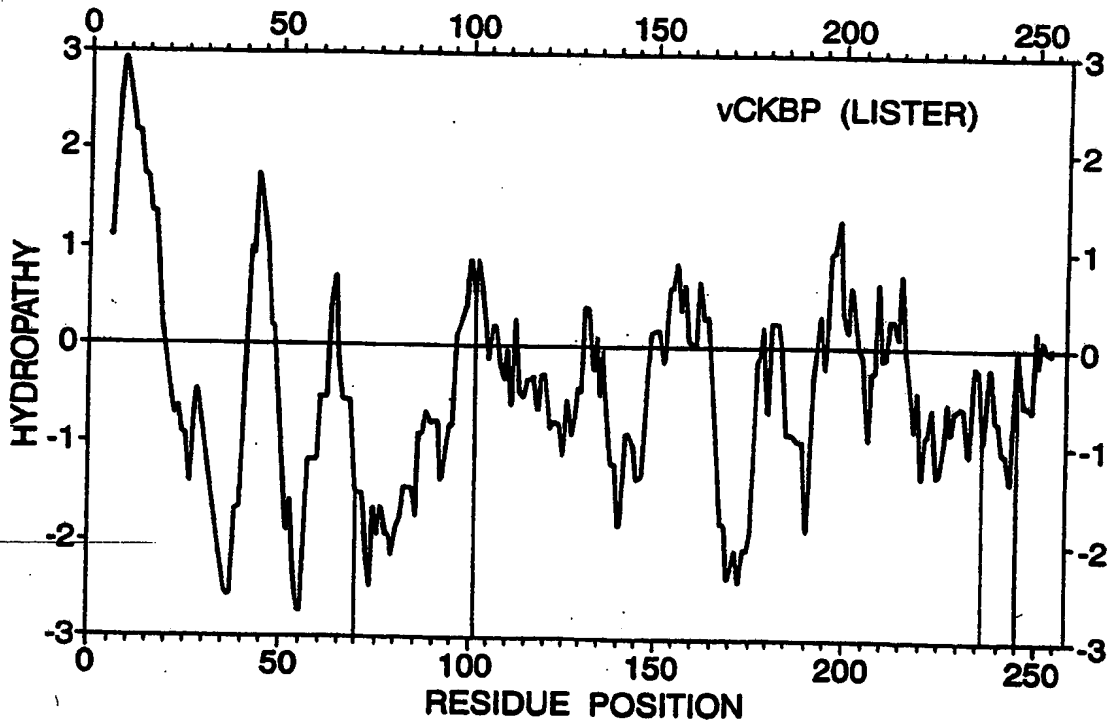
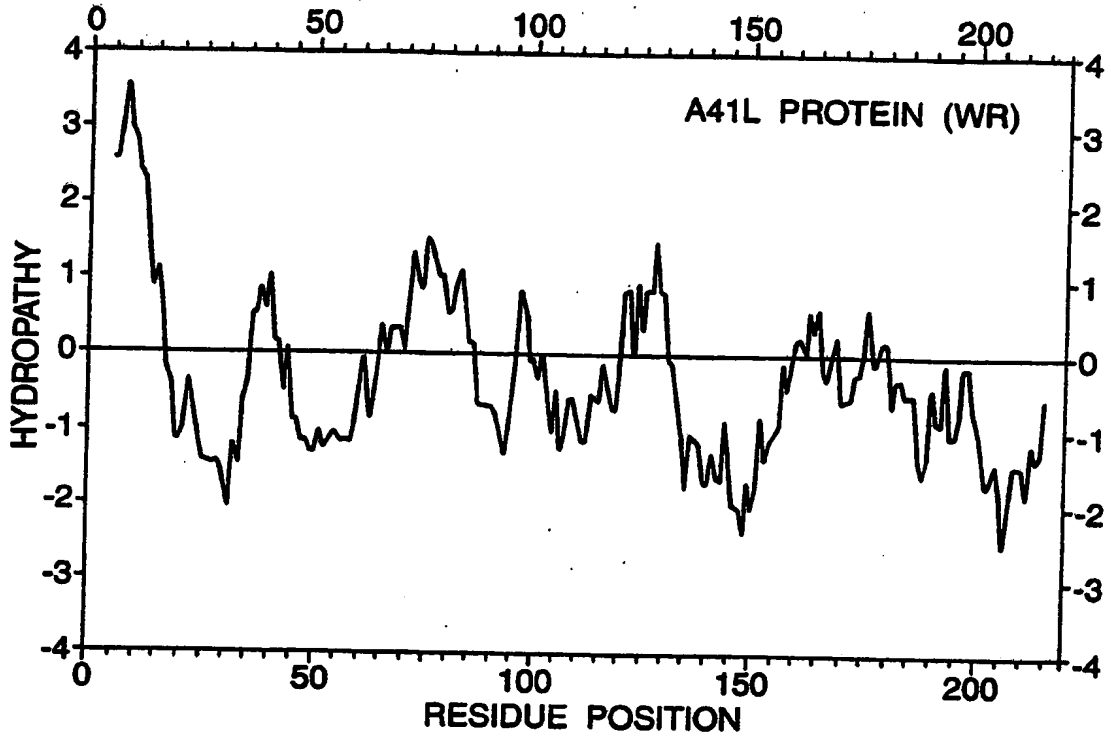


Fig.3.

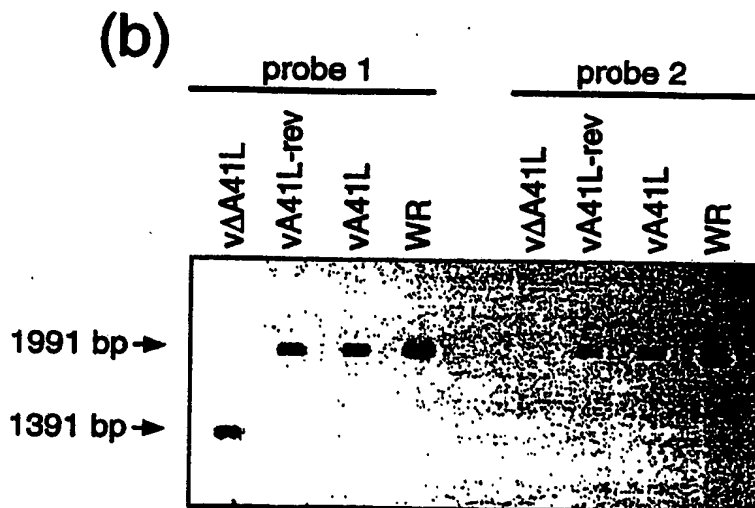
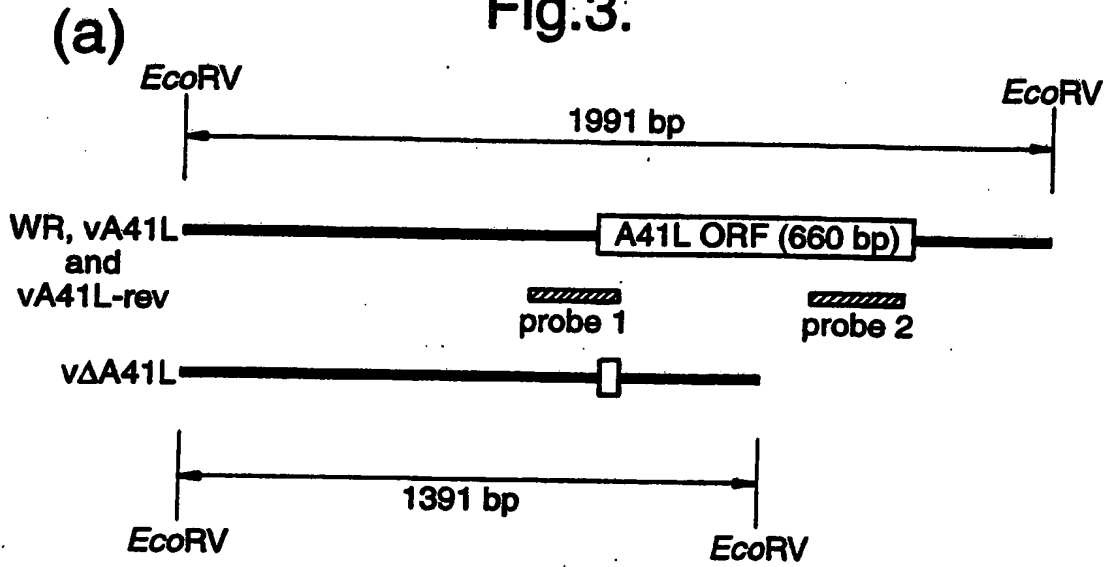


Fig.4.

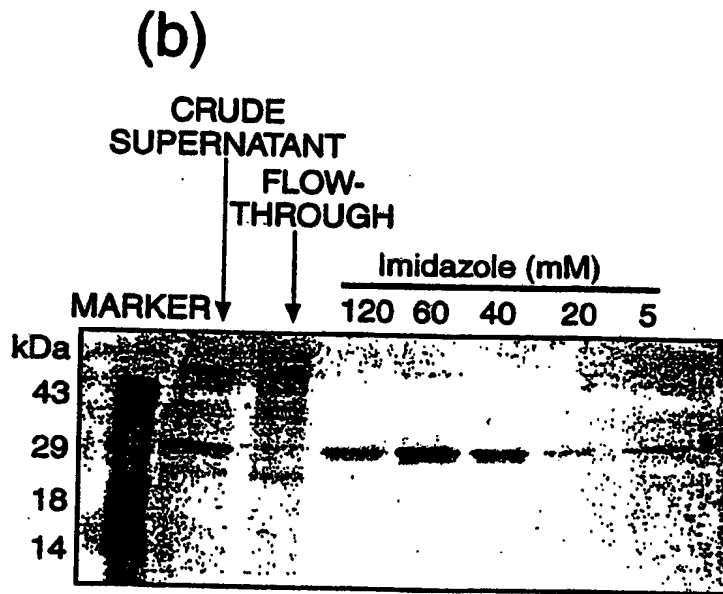
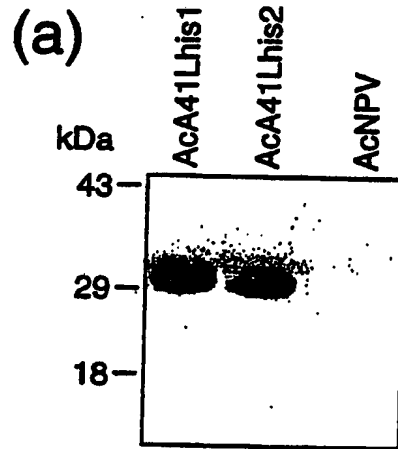
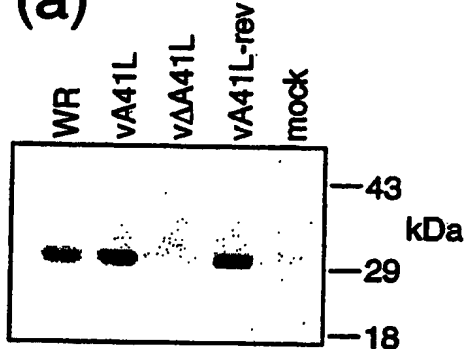
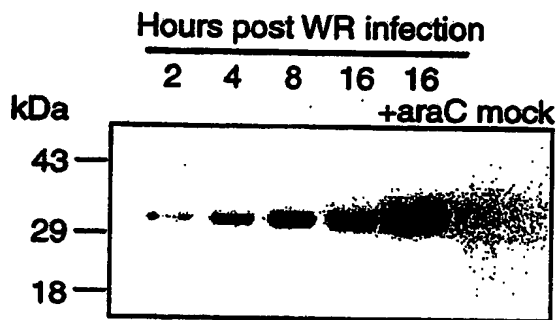


Fig.5.

(a)



(b)



(c)

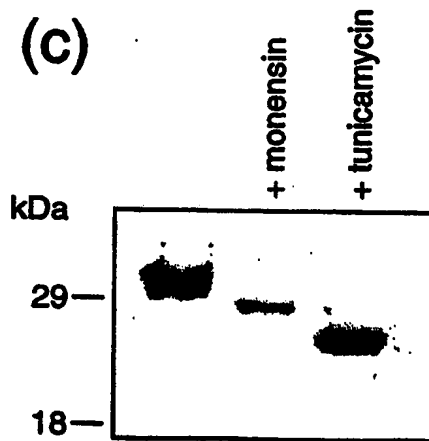


Fig.6.

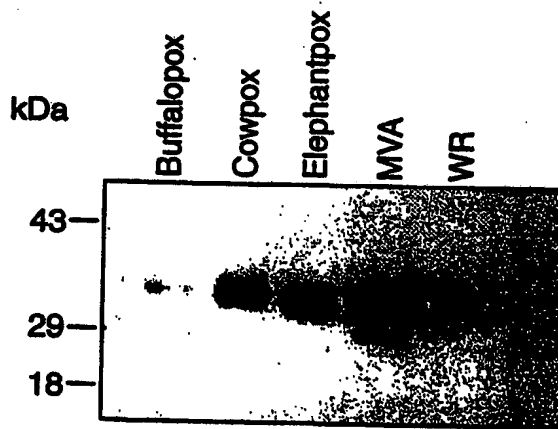
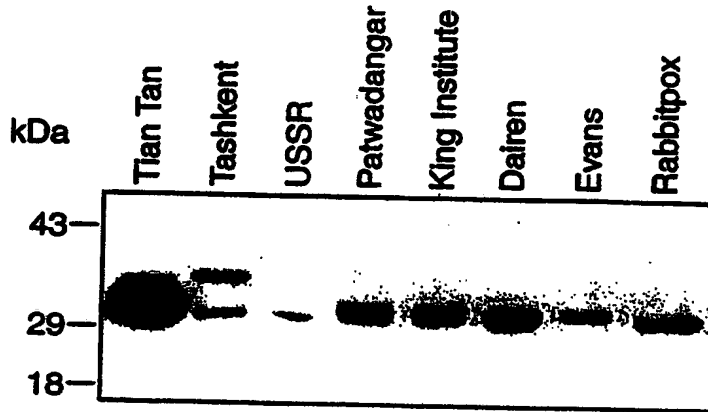
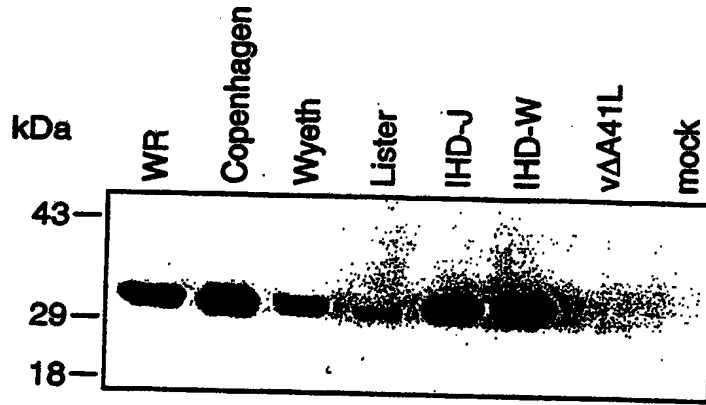


Fig.7(a).

CXC CHEMOKINES

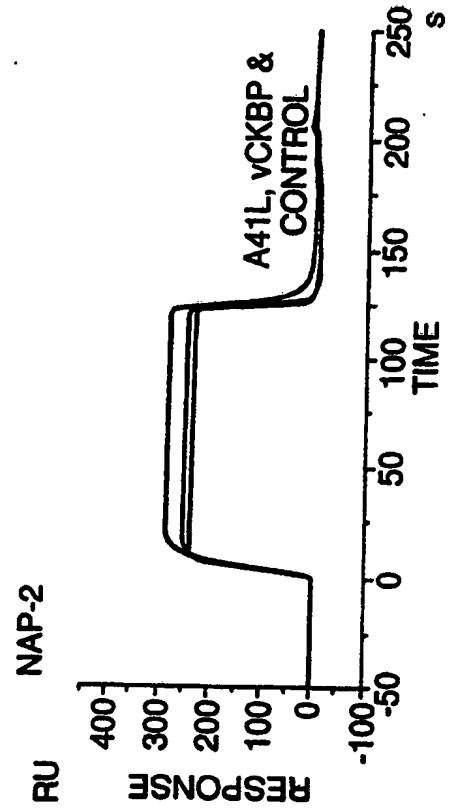
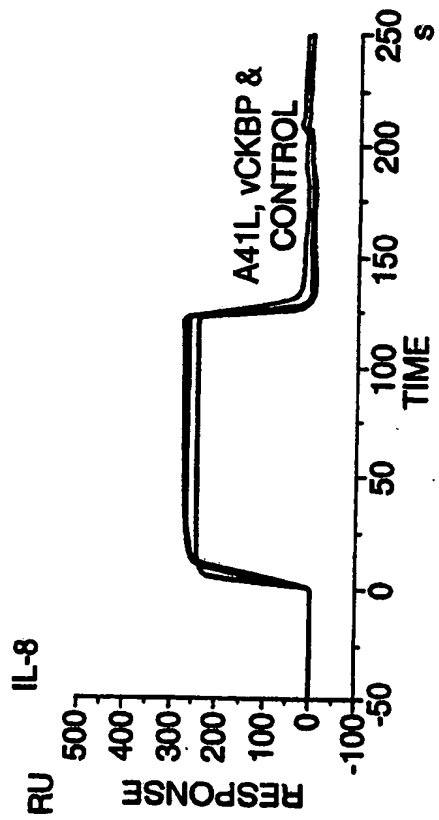
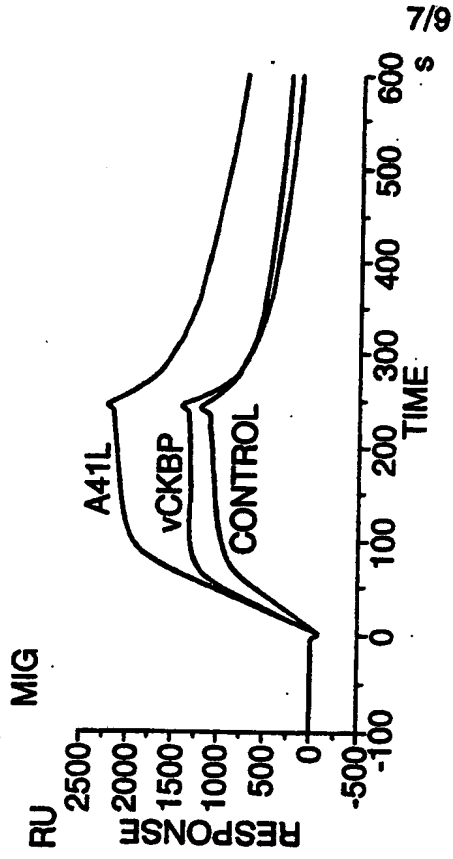
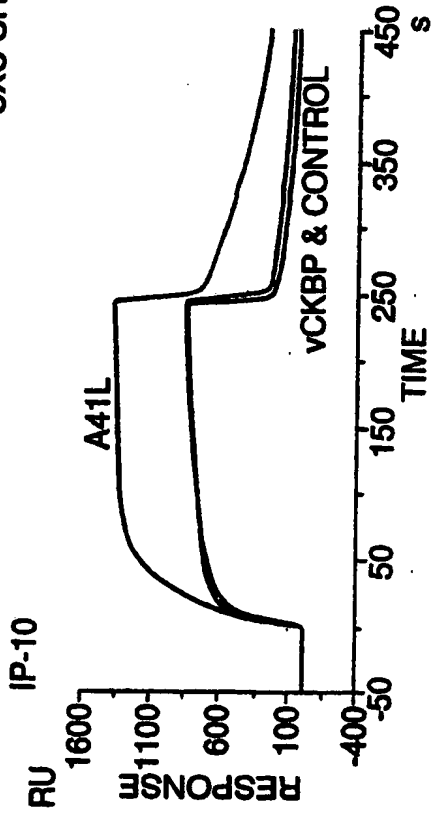


Fig.7(b).

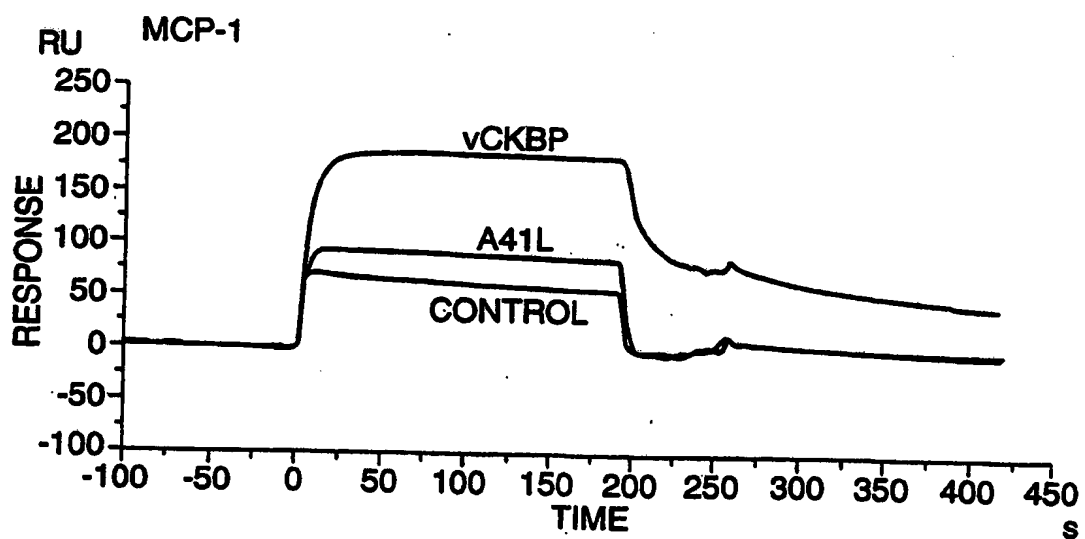
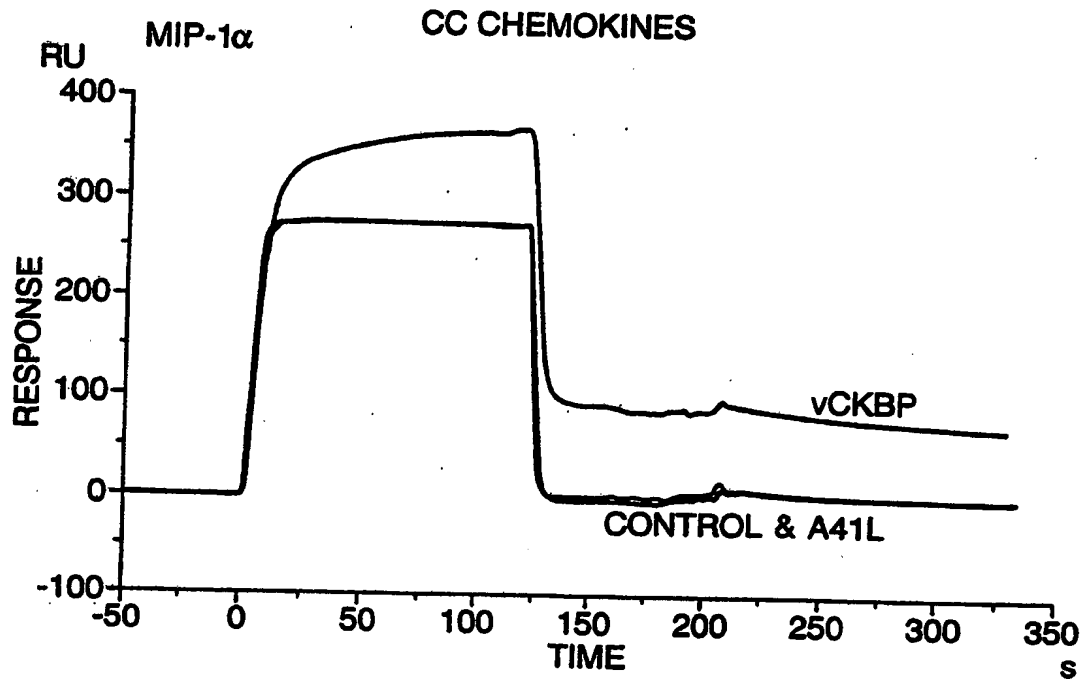


Fig.8.

1 AAACTTGACA TTAGCATT~~TT~~ ATTCTTATTA CAAATATAA AATAAAATAT
 51 ACAATCCAAT ACTCACATAA TCCA~~ACT~~CAC TCGAACACTA TTTTTCCAAT
 101 TACGATAACA ATATTGCAGA ATGTACTCGT TAGTATTTGT TATTTTGATG
 151 TGTATACCAT TTAGTTTICA AACAGTGTAT GATGATAAAT CGGTATGCGA
 201 TTCTGACAAT AAAGAATATA TGGGAATAGA AGTTTATGTA GAAGCAACGC
 251 TAGACGAACC CCTCAGACAA ACAACGTGTG AATCCAAAAT CCATAAATAT
 301 GGTGCATCTG TATCAAACGG AGGATTAAAT ATTTCTGTTG ATCTATTAAA
 351 CTGTTTTCTT AATTTTCATA CAGTTGGTGT ATACACTAAT CGCGATACCG
 401 TATACGCGAA GTTTGCTAGT TTGGATCCAT GGACTACGGA ACCTATAAAT
 451 TCTATGACCC ATGACGATCT AGTAAAATTA ACAGAAGAAT GTATAGTGGA
 501 CATTTATTTA AAATGTGAAG TGGATAAAAC AAAGGATTTT ATGAAAAC~~TA~~
 551 ACGGTAATAG ATTAAAACCA AGAGACTTTA AAACTGTTCC TCCTTCTAAT
 601 GTAGGAAGCA TGATAGA~~ACT~~ ACAGTCTGAC TATTGCGTAA ACGATGTGAC
 651 TACATACGTC AAAATATACG ATGAGTGTGG AAACATTAAA CAGCATTCCA
 701 TTCCAACACT AAGAGATTAT TTACCACCA AGAATGGTCA ACCACGTAAA
 751 ATATTAAAGA AAAAATTTGA TAATTGTTAA TTGTTATTTT TATAAAAACA
 801 AGAACGGTAC GGCGATATTT ATTTTTTCT AAAACATCTA ACCGAAGTAG
 851 TGGTATGATA AAAATGTAGT GTAATTGTTA TATAGTGTA CACGAAT